Applications of Genetic Algorithms and Quantum Genetic Algorithms on theoretical fully-autonomous road systems

University of Birmingham



 $\begin{array}{c} {\rm Sam~Barrett} \\ {\rm sjb786@student.bham.ac.uk} \end{array}$

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Abstract

Introduction

Background

2.1 Genetic Algorithms

2.1.1 History

Genetic algorithms are optimisation techniques that employ the same rationale as classical Evolution as seen in nature.

Genetic Algorithms can trace their origins back to the late 1960s when they were first proposed by John Holland, though he then referred to them as Genetic Plans¹. Holland went on to write the first book on the subject titled Adaptation in Natural and Artificial Systems[1] in 1975. The field did not find much reception with Holland stating in the preface to the 1992 rerun:

"When this book was originally published I was very optimistic, envisioning extensive reviews and a kind of 'best seller' in the realm of monographs. Alas! That did not happen."

However, in the early nineties, Genetic algorithms surged in popularity along with the area of Artificially Intelligent planning as a whole leading to Holland republishing his book and solidifying his position as the field's founder.

2.1.2 Definition

In a general sense, optimisation techniques work to find the set of parameters \mathcal{P} that minimise an objective function \mathcal{F} . Genetic algorithms approach this

 $^{^1}$ This distinction was made to emphasise the "centrality of computation in defining and implementing the plans" [1]

by representing these sets as individuals in a population, P. Over the course of multiple generations, the best solutions are determined and promoted until termination criteria are met or the maximum number of generations is reached.

As our candidates are essentially a collection of parameters to the function we are trying to optimise, we can extend our metaphor further by mapping each element of a individual to a *gene* in a individual's genome.

The representation we use in a GA is problem specific. Often we have to provide functions to facilitate the mapping between the problem specific set of possible solutions and the encoded genotype space in which we optimise. The most basic representation being a string of binary numbers.

Genetic algorithms are both probabilistically optimal and probabilistically complete [2] meaning that: given infinite time, not only will the algorithm find a solution, (if one exists), it will find the optimal solution from the set of all possible solutions, \mathcal{P}^* .

Algorithm 1: Modern Generic Genetic Algorithm

```
Result: Best Solution, p_{\texttt{best}}
Generate initial population, P_0 of size n;
Evaluate fitness of each individual in P_0, \{F(p_{0,1},\ldots,p_{0,n})\};
while termination criteria are not met do

Selection: Select individuals from P_t based on their fitness;
Variation: Apply variation operators to parents from P_t to produce offspring;
Evaluation: Evaluate the fitness of the newly bred individuals;
Reproduction: Generate a new population P_{t+1} using individuals from P_t as well as the newly bred candidates.;
t++
end
return p_{\texttt{best}}
```

As you can see from Figure 2.1 and Algorithm 1 the overall shape of GAs has not changed substantially over the course of the past 50 years. Being comprised of a series of operations that a starting population is piped through until criteria are met.

2.1.3 Selection

The selection procedure is the process by which the next generation of individuals is created from the current population. Individuals are selected relative to their fitness as determined by the Objective (fitness) function \mathcal{F} .

```
Set t = 0 and initialize B by selecting M structures at random from a_1 to
     form \mathfrak{B}(0) = (A_1(0), \ldots, A_M(0)).
 2.1 Observe and store the performances (\mu_1(0), \ldots, \mu_M(0)) to form \mathcal{U}(0).
  2.2 Calculate \hat{\mu}(0) = \sum_{h=1}^{M} \mu_h(0)/M.
        ▶ 2.3 Observe the performance \mu_E(A'(t)).
           2.4 Update \hat{\mu}(t) by calculating \hat{\mu}(t) + \mu_B(A'(t))/M - \mu_{i(t)}(t)/M.
           2.5 Update V(t) by replacing \mu_{j(t)}(t) with \mu_E(A'(t)).
     Increment t by 1.
     Define the random variable Rand_t on \theta_M = \{1, \ldots, M\} by assigning the
     probability \mu_h(t)/\hat{\mu}(t) to h \in \mathcal{G}_M. Make one trial of Rand_t and designate
     the outcome i(t).
 5.1 Apply simple crossover (as defined in section 6.2 and extended in section
     6.3) to A_{i(t)}(t) and A_{i'(t)}(t) with probability P_c, where A_{i'(t)}(t) is determined
      by a second trial of Randt. Select one of the two resultants at random
     (equilikely) and designate it {}^{1}A(t) (where the order of attributes in the
     resultant is that of A_{i(t)}(t).
 5.2 Apply simple inversion (as defined in section 6.3) with probability P_L
     yielding {}^{2}A(t).
 5.3 Apply mutation (as defined in section 6.3) to {}^{2}A(t) with probability
     c_t \cdot {}^{1}P_M, yielding A'(t).
 6.1 Assign probability 1/M to each h \in \mathcal{G}_M and make a random trial accord-
     ingly; designate the outcome j(t).
←6.2 Update \mathfrak{B}(t) by replacing A_{j(t)}(t) with A'(t).
```

Figure 2.1: GA algorithm outlined in Holland's Original Book[1]

Some methods select only the best solutions by fitness. Others employ a more stochastic approach such as the Monte Carlo method[3] to increase diversity and reduce complexity.

2.1.4 Variation

Variation in a GA is the process of altering the genome individuals to further explore the search space via stochastic local search. We perform this using two distinct sub-operations: Mutation and Crossover.

Mutation

In mutation we alter each gene with a set probability p_m known as the mutation rate. A standard value for a mutation rate is $\frac{1}{L}$ but it can fall anywhere in the range $p_m \in [\frac{1}{L}, \frac{1}{2}]$ where L is the length of the genome. In binary coded GAs we alter a given gene by flipping it's value. In real-coded GAs mutation operators include:

- Uniform Mutation
- Non-Uniform Mutation
- Gaussian Mutation

A low value for p_m a new individual which can be shown to be *close* to it's parents in the search space relative to their $Hamming\ distance^2$ if using a Binary coded GA. In real-coded GAs they can be shown to be close by the Euclidean distance between them.

Crossover

Crossover is a binary operation, taking two randomly selected parents from the population P_t with a probability $p_c \in [0, 1]$

There are two major forms of crossover for binary coded GAs: n > 1 point crossover and Uniform crossover.

For real-coded GAs you instead have a selection of Crossover operators including:

- Flat crossover
- Simple crossover

²Hamming Distance: The metric for comparing two binary data strings. The Hamming distance between two strings is the number of bit position in which they differ.

- Whole arithmetic crossover
- Local arithmetic crossover
- Single arithmetic crossover
- BLX- α crossover

2.1.5 Evaluation

After developing potential new individuals, the fitness of these new individuals is calculated using the objective function, \mathcal{F}

2.1.6 Reproduction

Here the next generation of individuals is constructed. There are multiple potential methods employed here with the simplest being to just replace the lest fit individuals with additional copies of the fitter individuals, be they pre-existing or newly generated.

In this stage various heuristics or alternative strategies can be implemented to speed up or slow down the convergence rate. Whilst their fitness may be low, having a diverse population allows for more of the search space to be explored. If the algorithm converges too quickly it may get *stuck* in local minima.

2.2 Fully Autonomous Road Networks

Fully autonomous road networks are by no means a novel concept. They have appeared in fiction since the mid 20th century and have been a real possibility for the past decade. In a fully autonomous road network (FARN), all vehicles are self-operating with their routing either being self-serving or determined by a larger system managing routes for all nearby agents.

 FARN s have the possibility of improving travel in a number of different areas.

They have the potential of greatly reducing travel times by efficiently routing all vehicles with the goal of reducing the net travel time. Such a system would also be able to respond much faster than human drivers, allowing for large increases in permissible vehicle velocities thus, further increasing efficiency.

By extension of their higher efficiency, FARNs will also have a marked effect on vehicular energy consumption. The potential effects are summarised

nicely by Vahidi and Sciarretta[4] where they show the use of vehicle automation could cause anywhere from a halving of "energy use and greenhouse gas emission" in an "optimistic scenario" to doubling them depending on the effects that present themselves. The increase in highway speed whilst increasing travel efficiency is predicted by Brown et al.[5] and Wadud et al.[6] as increasing energy use by anywhere from 5% to 30%

2.3 Quantum Genetic Algorithms

2.4 Alternative Technologies

Literature Review

3.1 Classical GAs

3.2 Fully Autonomous Road Networks

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3.3 Quantum GAs

3.3.1 Quantum Computing

Classical Approach

- 4.1 Approach
- 4.2 Implementation
- 4.2.1 Language Choice
- 4.3 Results

Quantum Approach

- 5.1 Approach
- 5.2 Implementation
- 5.3 Results

Evaluation

Conclusion

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